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1/15

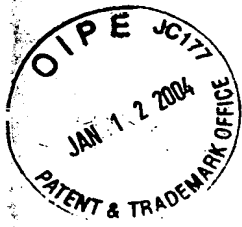


Figure 1a

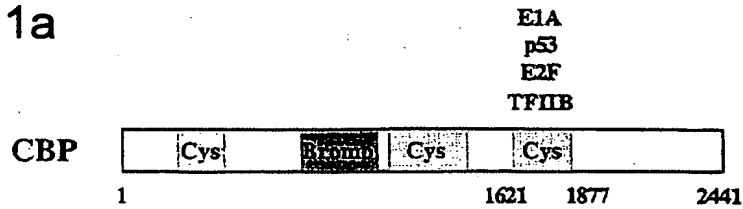
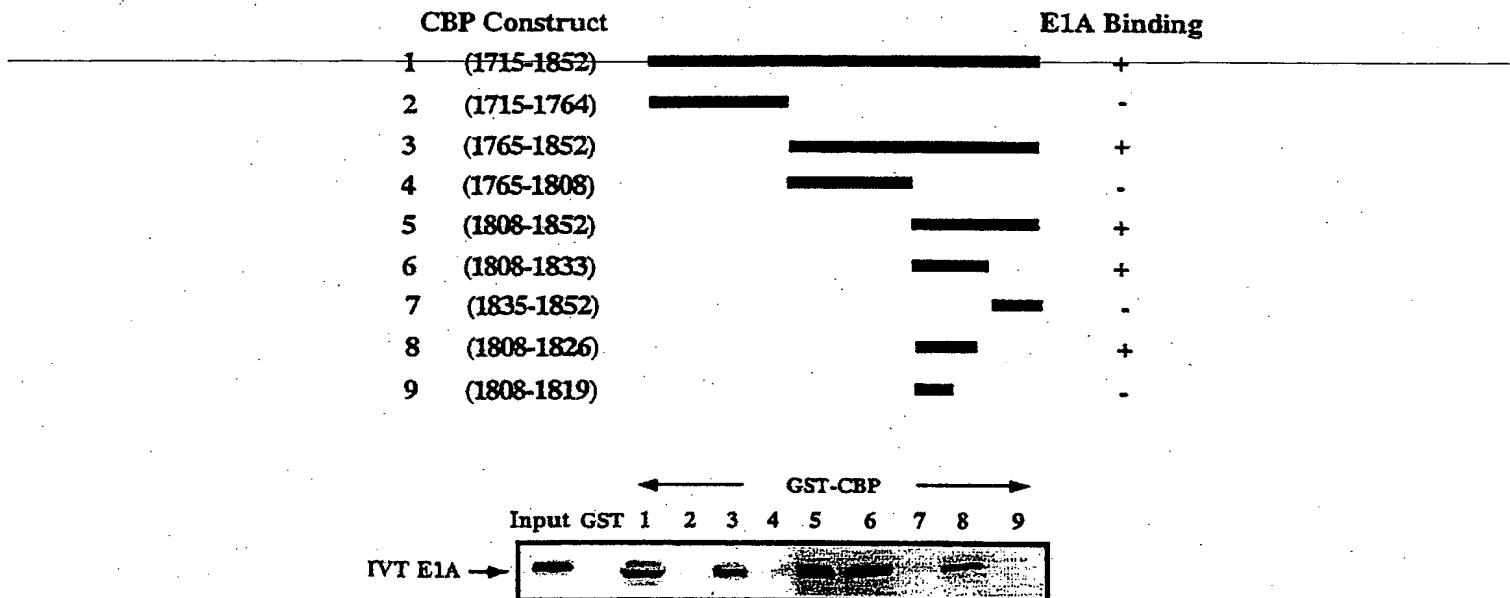


Figure 1b



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2/15

Figure 1c

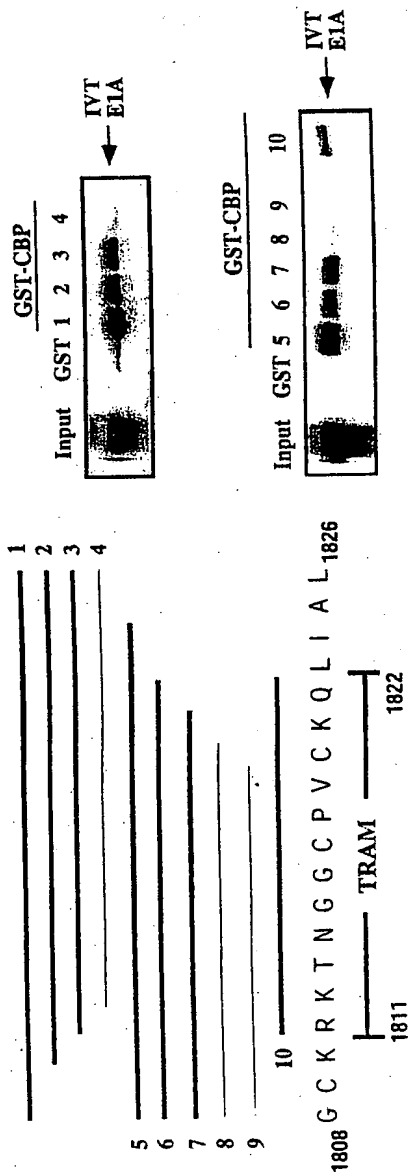
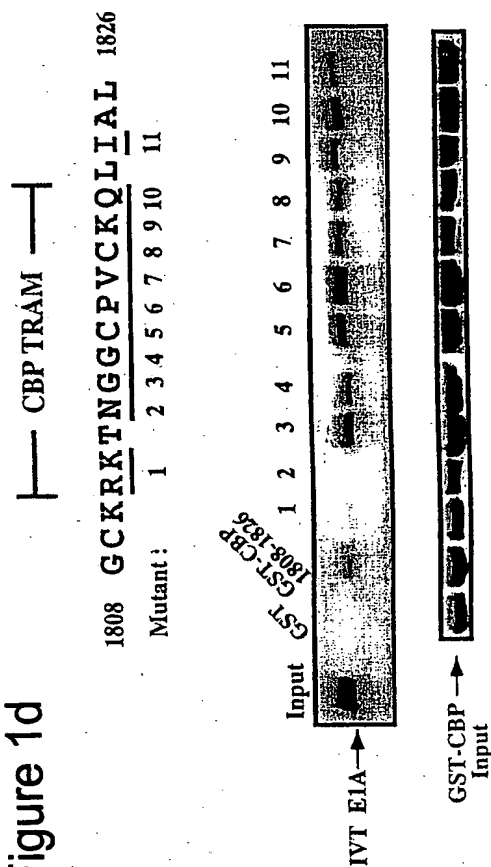


Figure 1d



3/15

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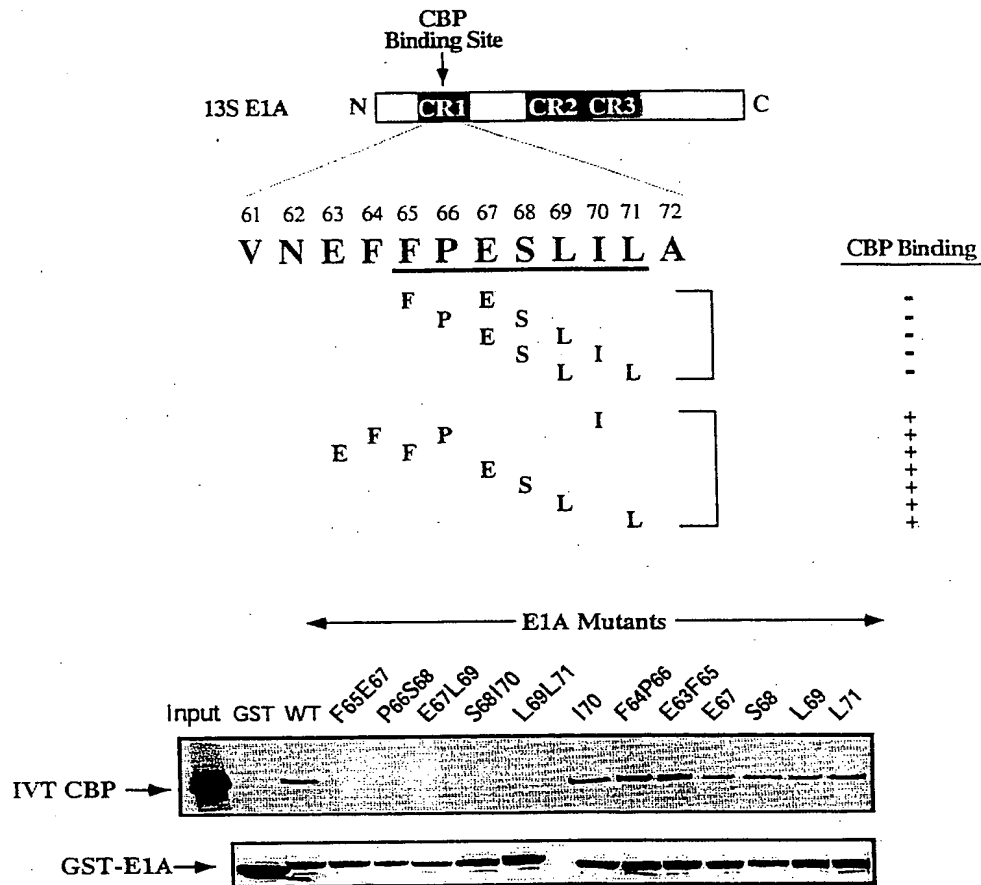
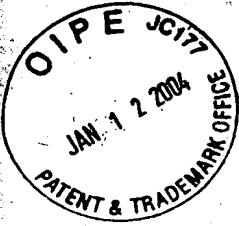
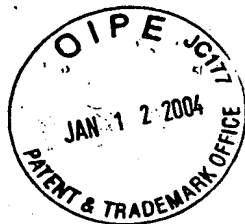


Figure 2a



4/15

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Figure 2b

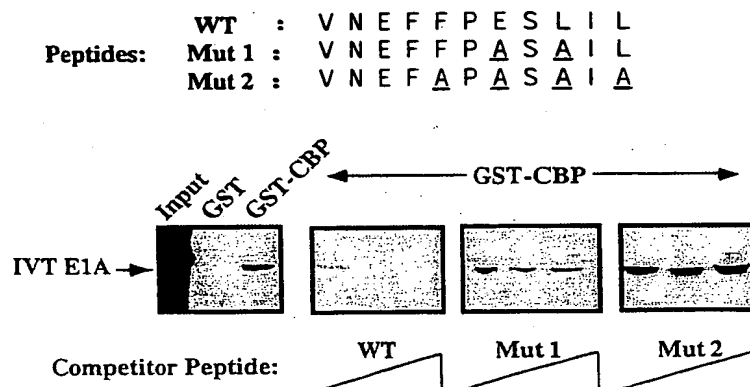


Figure 2c

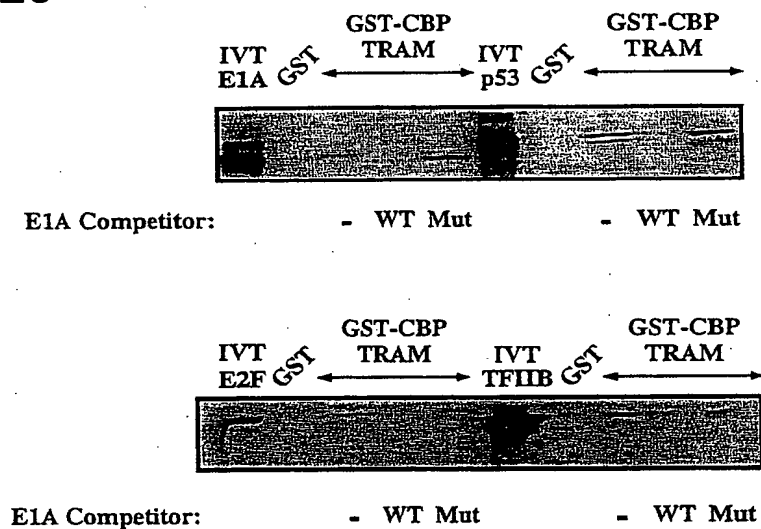


Figure 2d

TRAM	Interaction Motif	E1A	V N E F <u>F</u> P E S L I L A A
p53		S Q E T <u>F</u> S D L W K L L P	
E2F		F D C D <u>F</u> G D L T P L D F	

Consensus: F X $\frac{E}{D}$ X X X L



5/15

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Figure 3a

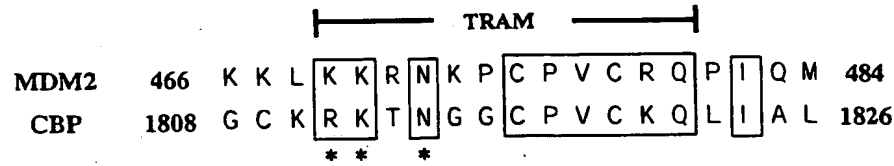


Figure 3b

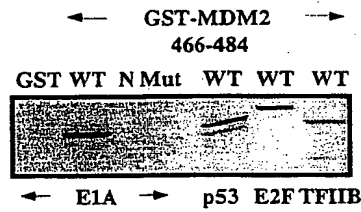


Figure 3c

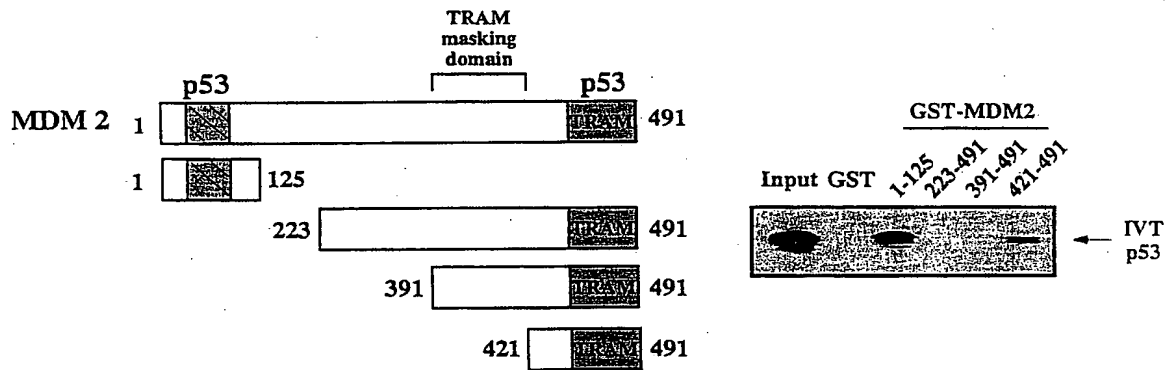
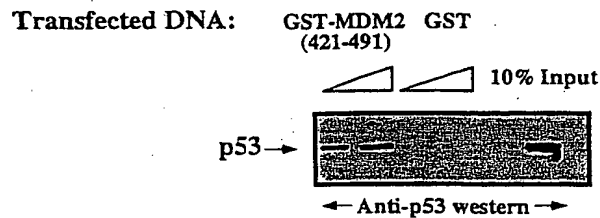
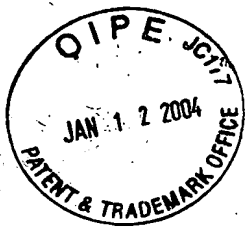


Figure 3d





6/15

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Figure 4a

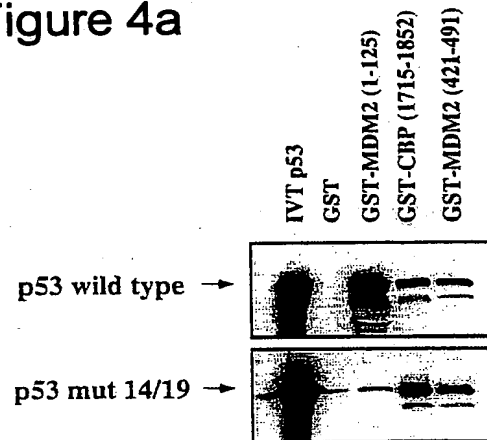


Figure 4b

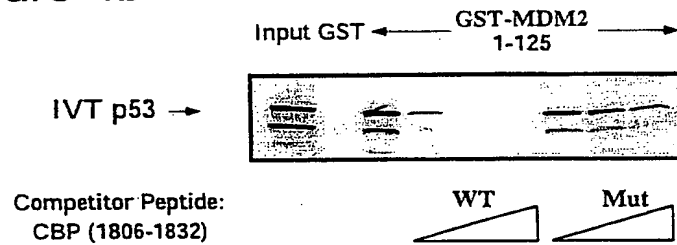


Figure 4c

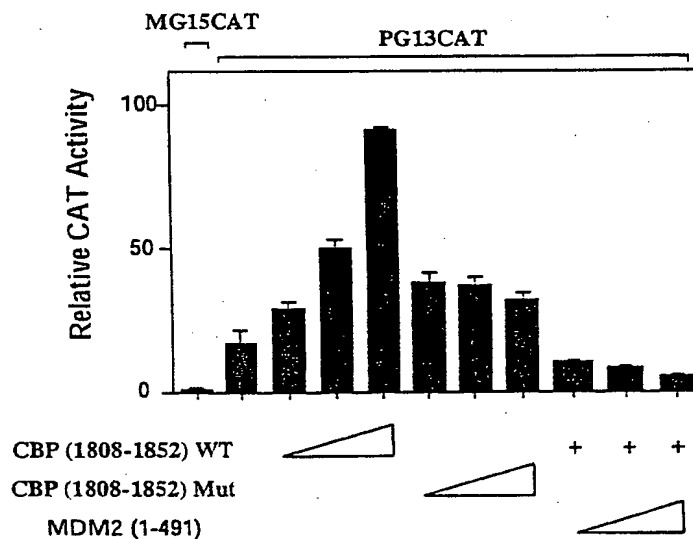
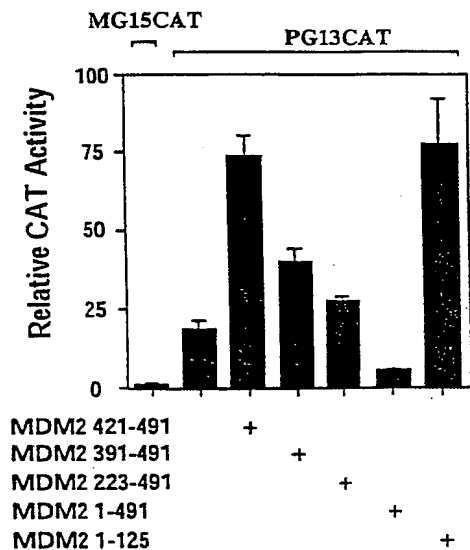


Figure 4d





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7/15

Figure 5a

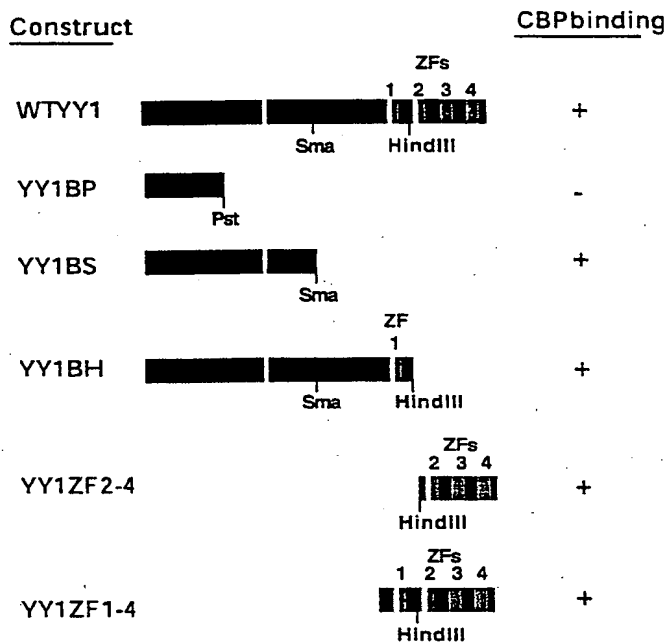


Figure 5b

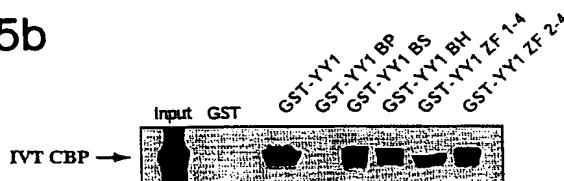
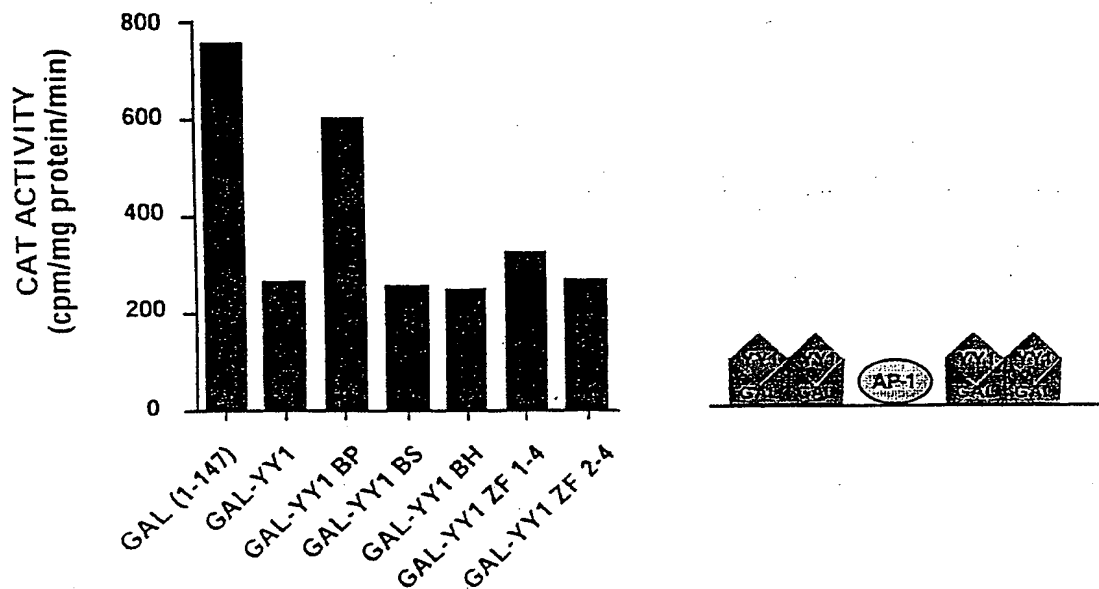


Figure 5c



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TRIMs

└

— CBP TRAM —

Input GST GST-CBP 1808-1826 1 2 3 4 5 6 7 8 9 10 11

IVT E1A →

Input GST GST-CBP 1808-1826 GST 1 2 3 4 5 6 7 8 9 10 11

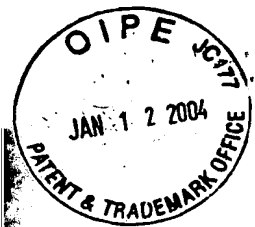
IVT p53 →

Input GST GST-CBP 1808-1826 1 2 3 4 5 6 7 8 9 10 11

IVT TFIIIB →

Input GST GST-CBP 1808-1826 1 2 3 4 5 6 7 8 9 10 11

IVT YY1 →



9/15

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Figure 7a

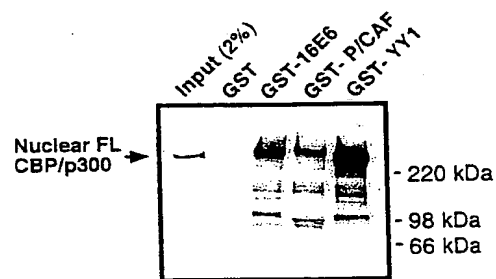


Figure 7b

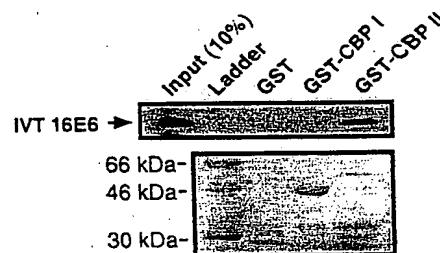


Figure 7c

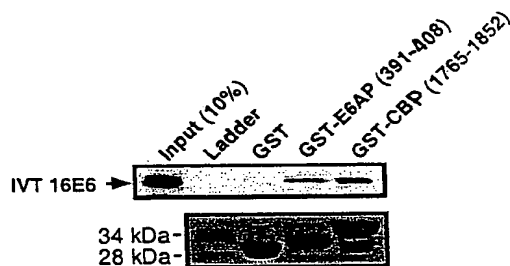
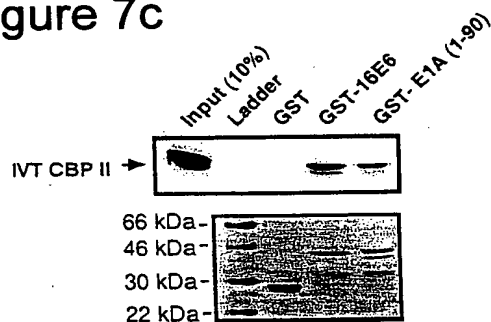
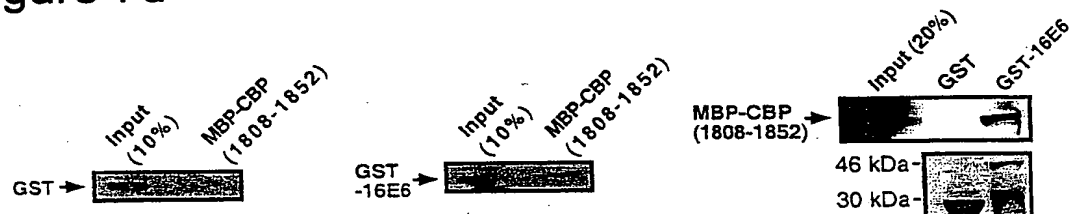


Figure 7d

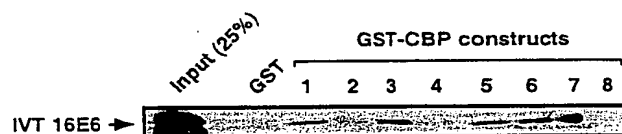




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10/15

	CBP constructs	E6 binding
1	1715 1870	+
2	1715 1764	-
3	1765 1852	+
4	1765 1808	-
5	1808 1852	+
6	1808 1833	+
7	1808 1826	+
8	1808 1819	-



CBP (1808-1826): GCKRKTNGGCPVCKQLIAL
 p300 (1770-1788): GCKRKTNGGCPICKQLIAL
 *****+*****



Figure 8



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11/15

Figure 9a

16 E6

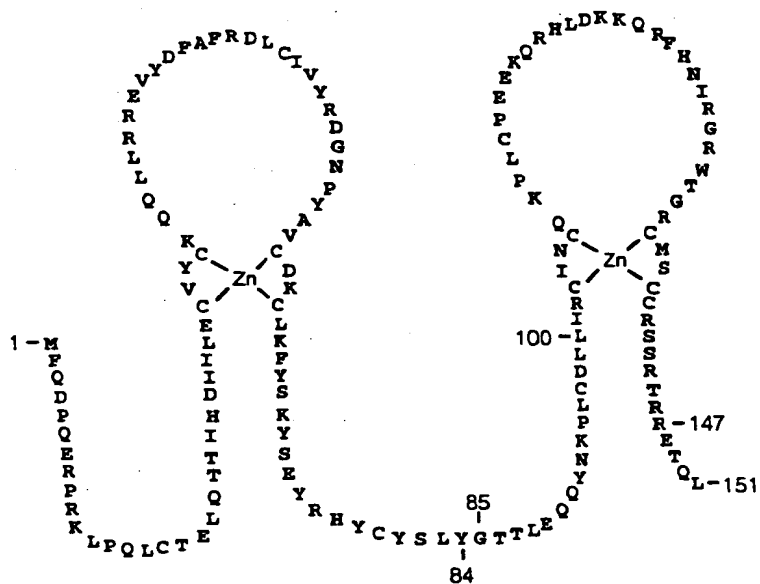
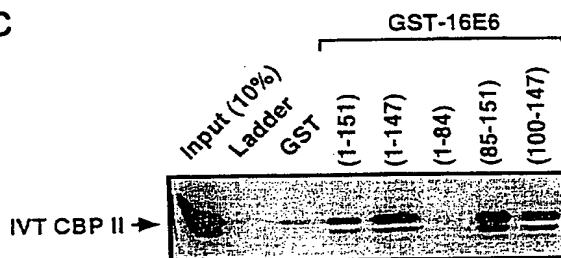


Figure 9b

16 E6 constructs	CBP binding
1-151	+
1-147	+
1-84	-
85-151	+
100-147	+

Figure 9c





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Figure 10a

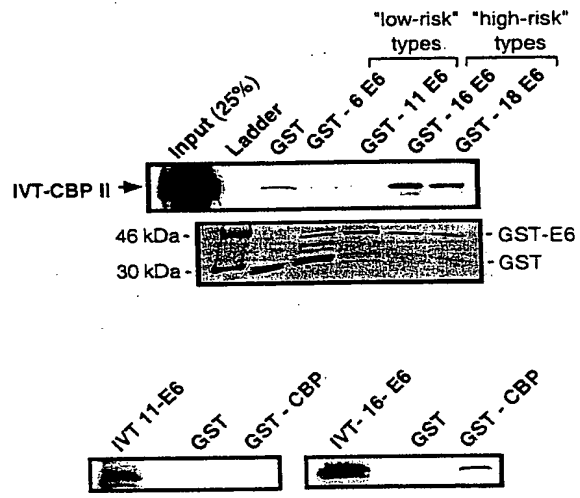
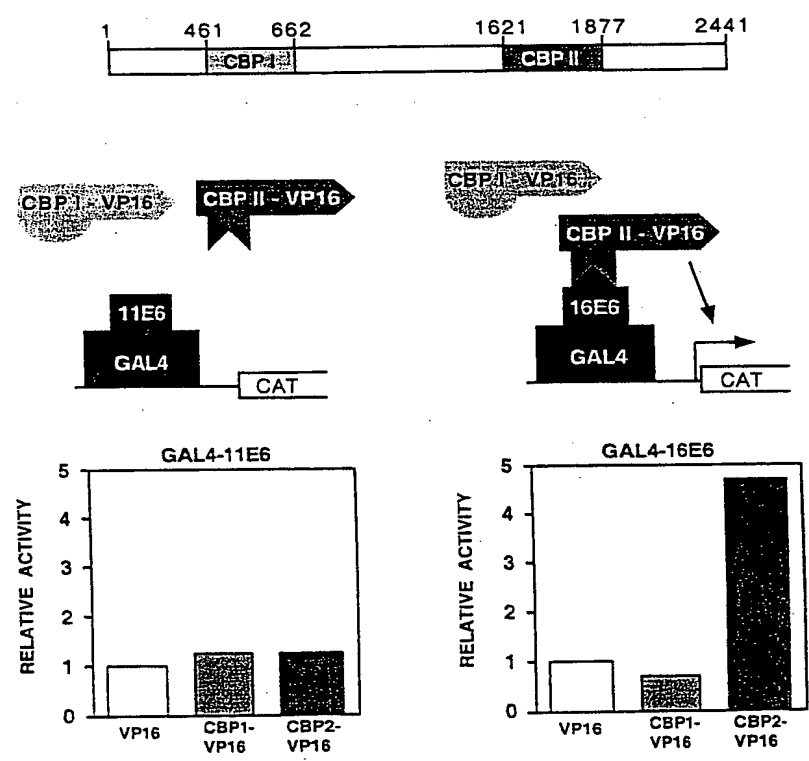


Figure 10b





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Figure 11a

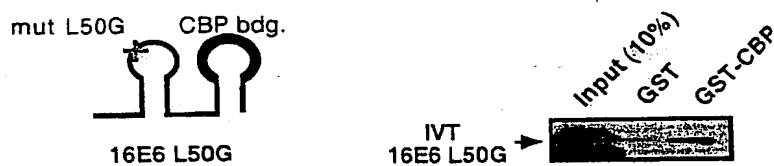


Figure 11b

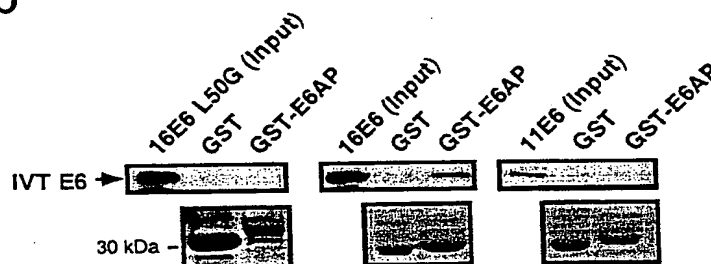
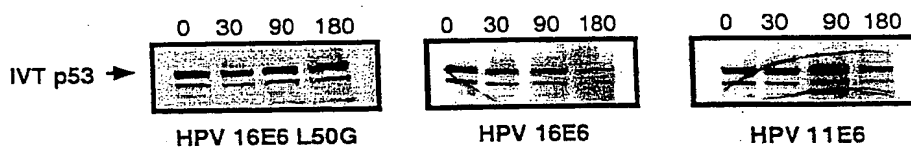


Figure 11c





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14/15

Figure 12a

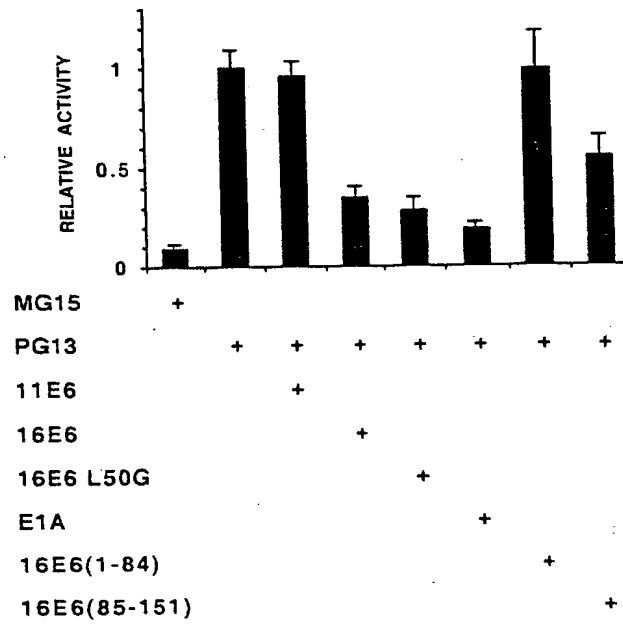
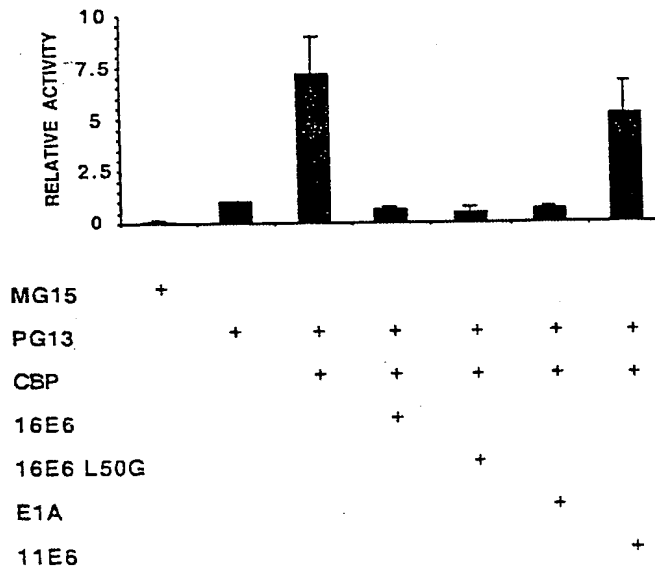


Figure 12b





15/15

G	S	V	D	L	E	P	G	V	Q	K	L	x	x	x	
GGA	TCC	GTC	GAC	CTC	GAG	CCC	GGG	CTG	CAG	AAG	CTT	GAT	TGA	Tta	gct t 3'
cct agG	CAG	CTG	GAG	CTC	GAG	CCC	GGG	GTC	GTC	TTC	GAA	CTA	ACT	AAT	GGA a 5'
BamH1				Xho1		Sma1		Pst1		HindIII		stop (x3)			[core HindIII]

Figure 13